

Supplementary Materials

Synthesis, polymorphism and thermal decomposition process of $(n\text{-C}_4\text{H}_9)_4\text{NRE}(\text{BH}_4)_4$ for $RE = \text{Ho, Tm and Yb}$

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1. PXRD Patterns, Rietveld Refinement and Structural Information

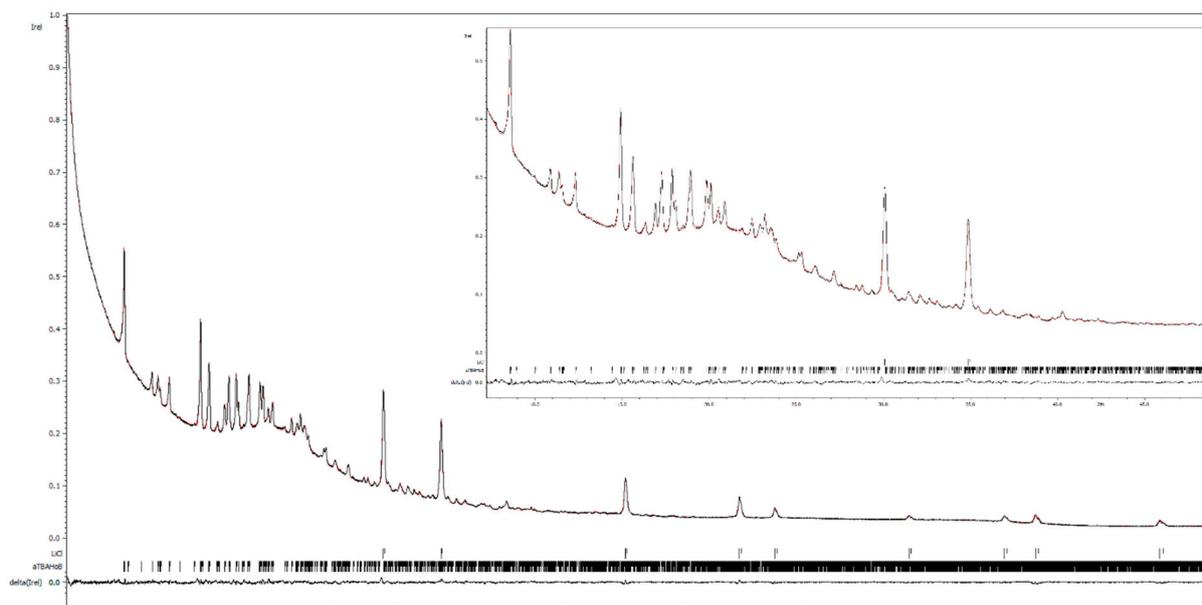


Figure S1. Rietveld refinement for as-milled sample **Ho**. Black curve represents experimental data, red curve – calculated profile. The position of the Bragg reflections has been marked and the difference curve (between the experimental and calculated profiles) are plotted at the bottom. Inset: the low angle region. GOF = 1.73; Rp = 0.73; wRp = 1.06. Wavelength: Cu ($K_{\alpha 1}$ and $K_{\alpha 2}$). Bragg reflections marked, from top to bottom, for: LiCl and α -TBAHoB.

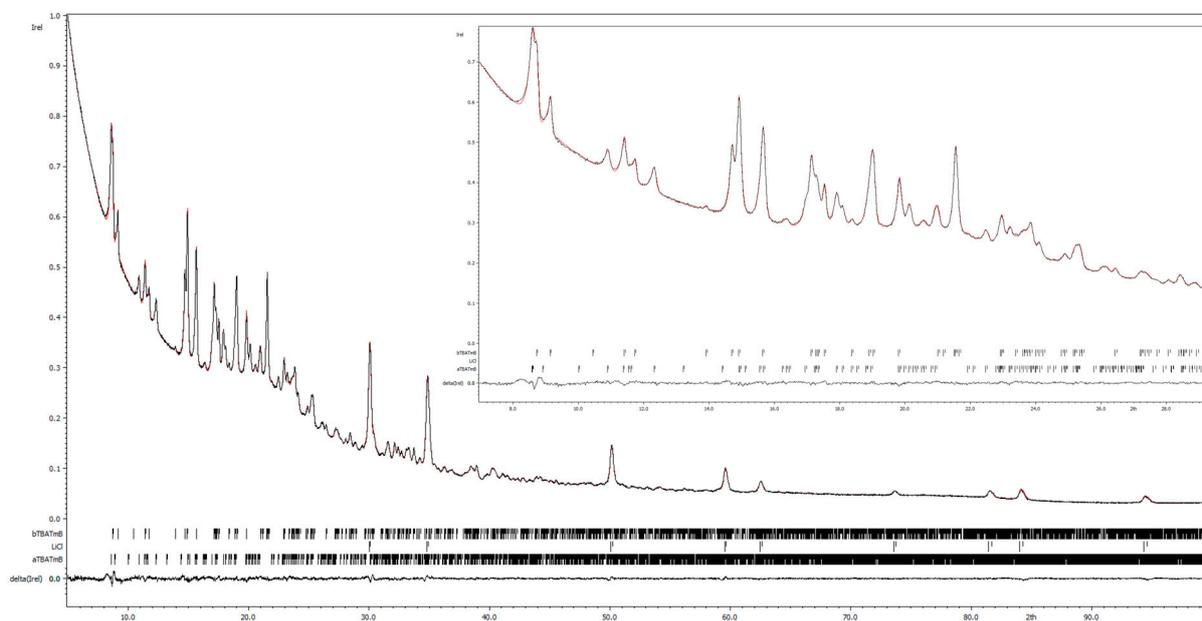


Figure S2. Rietveld refinement for as-milled sample **Tm**. Black curve represents experimental data, red curve – calculated profile. The position of the Bragg reflections has been marked and the difference curve (between the experimental and calculated profiles) are plotted at the bottom. Inset: the low angle region. GOF = 1.64; Rp = 0.69; wRp = 0.97. Wavelength: Cu ($K_{\alpha 1}$ and $K_{\alpha 2}$). Bragg reflections marked, from top to bottom, for: β -TBATmB, LiCl and α -TBATmB.

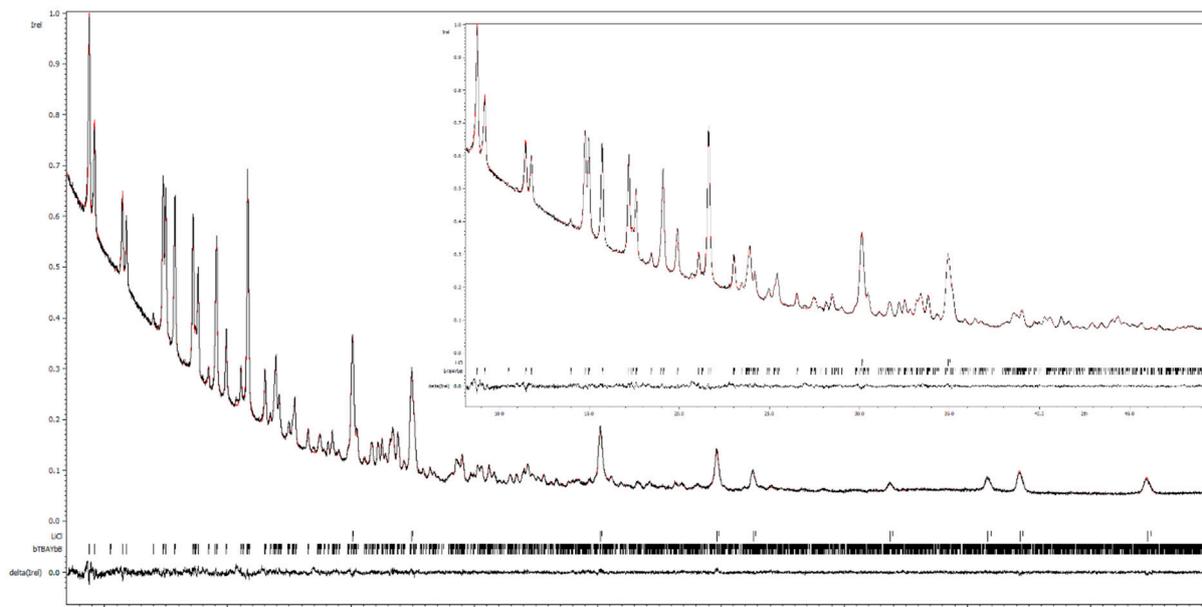


Figure S3. Rietveld refinement for as-milled sample **Yb**. Black curve represents experimental data, red curve – calculated profile. The position of the Bragg reflections has been marked and the difference curve (between the experimental and calculated profiles) are plotted at the bottom. Inset: the low angle region. GOF = 0.03; Rp = 1.29; wRp = 1.71. Wavelength: Cu ($K_{\alpha 1}$ and $K_{\alpha 2}$). Bragg reflections marked, from top to bottom, for: LiCl and β -TBA**Yb**B.

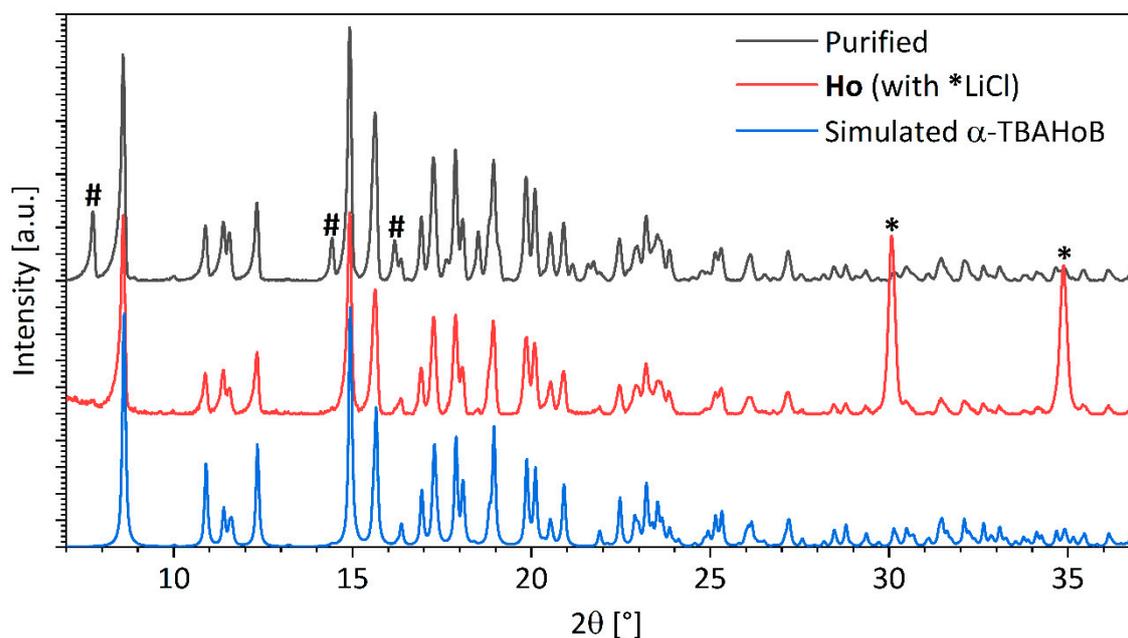


Figure S4. PXRD patterns with subtracted backgrounds for as-milled **Ho** sample (containing LiCl, marked *), purified **Ho** (with unknown impurities, marked #) and simulated pattern of α -TBA**Ho**B (from cif, crystal structure obtained from Rietveld refinement).

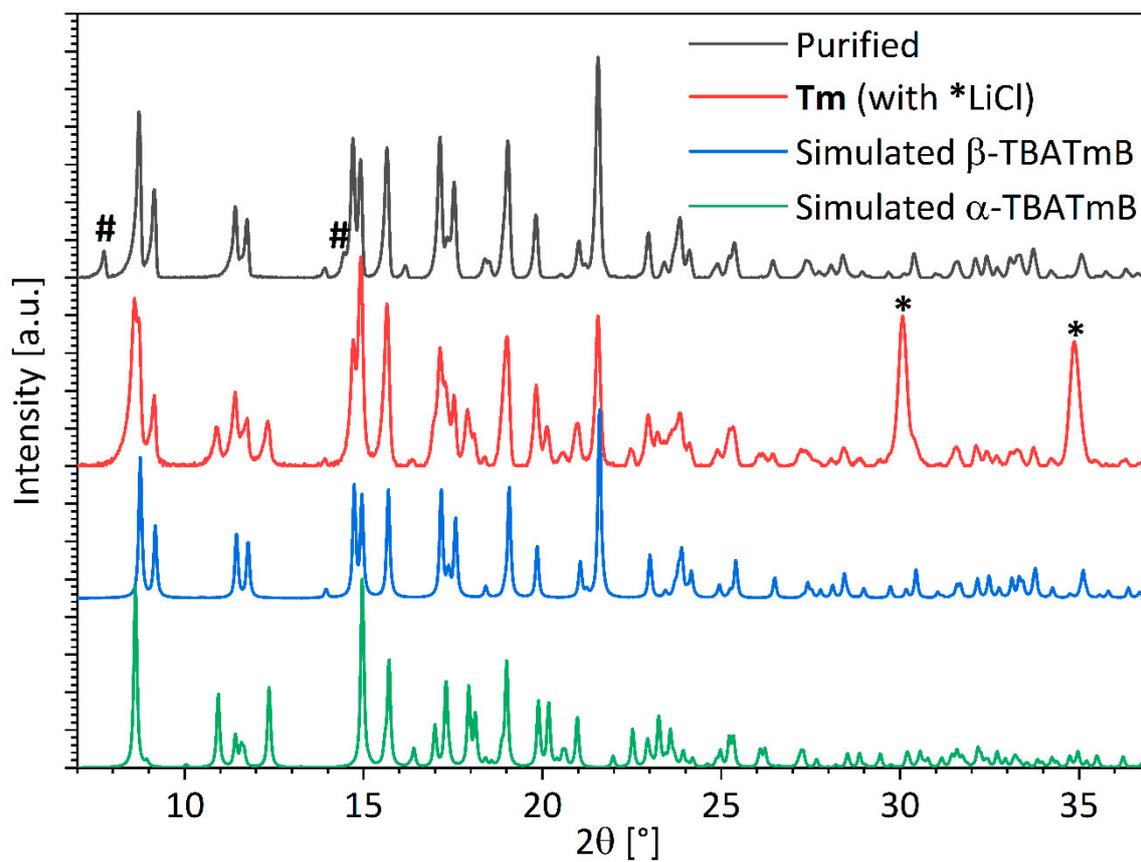


Figure S5. PXRD patterns with subtracted backgrounds for as-milled **Tm** sample (containing LiCl, marked *), purified **Tm** (with unknown impurities, marked #) and simulated patterns of α -TBATmB and β -TBATmB (from cifs, crystal structures obtained from Rietveld refinement).

Table S1. supplementary to **Table 2.** Unit cell dimensions for TBAREB, RE = Y, Ho, Tm, Yb, Sc. 100/200/300 K obtained from SC-XRD, RT from PXRD.

Compd.	α -TBAYB [41]		α -TBAHoB		α -TBA TmB	β -TBAHoB			β -TBATmB				β -TBAYbB		β -TBAScB [42]
RE ³⁺ r ¹ [Å]	0.900		0.901		0.880	0.901			0.880				0.868		0.745
spc. group	P2 ₁ /c		P2 ₁ /c		P2 ₁ /c	Pnna			Pnna				Pnna		Pnna
T [K]	100	RT	100	RT	RT	100	200	300	100	200	300	RT	100	RT	RT
a [Å]	11.0453(5)	11.4181(10)	11.039(3)	11.4218(9)	11.4063(18)	18.5597(8)	18.9387(15)	19.3238(12)	18.5303(5)	18.9730(6)	19.3409(16)	19.287(3)	18.5673(9)	19.2235(10)	19.1399(10)
b [Å]	20.0099(9)	20.510(3)	19.999(2)	20.553(2)	20.545(4)	11.9188(3)	11.9783(8)	12.0529(5)	11.8902(4)	11.9295(4)	12.0279(11)	12.0317(17)	11.8735(5)	11.9943(6)	11.8849(6)
c [Å]	14.7204(8)	15.2811(19)	14.708(4)	15.3049(17)	15.319(3)	11.7871(4)	11.8153(9)	11.8820(6)	11.7514(3)	11.7754(4)	11.8749(12)	11.8591(19)	11.7096(5)	11.8244(6)	11.7325(6)
β [°]	127.980(5)	129.464(8)	128.02(4)	129.433(7)	129.423(12)	90	90	90	90	90	90	90	90	90	90
V [Å ³]	2564.44	2762.77	2558.1(10)	2775.0(5)	2773.1(10)	2607.42(16)	2680.3(3)	2767.4(3)	2589.17(13)	2665.23(15)	2762.5(4)	2751.9(7)	2581.5(2)	2726.4(2)	2668.9(2)
Z	4		4		4	4			4				4		4

¹ Effective ionic radius (6-coordinate, octahedral environment) from [54]

Compd.	α -TBAYB [41]		α -TBAHoB		α -TBA TmB
RE ³⁺ r ¹ [Å]	0.900		0.901		0.880
spc. group	P2 ₁ /c		P2 ₁ /c		P2 ₁ /c
T [K]	100	RT	100	RT	RT
a [Å]	11.0453(5)	11.4181(10)	11.039(3)	11.4218(9)	11.4063(18)
b [Å]	20.0099(9)	20.510(3)	19.999(2)	20.553(2)	20.545(4)
c [Å]	14.7204(8)	15.2811(19)	14.708(4)	15.3049(17)	15.319(3)
β [°]	127.980(5)	129.464(8)	128.02(4)	129.433(7)	129.423(12)
V [Å ³]	2564.44	2762.77	2558.1(10)	2775.0(5)	2773.1(10)
Z	4		4		4

Compd.	β -TBAHoB			β -TBATmB				β -TBAYbB		β -TBAScB [42]
RE ³⁺ r ¹ [Å]	0.901			0.880				0.868		0.745
spc. group	Pnna			Pnna				Pnna		Pnna
T [K]	100	200	300	100	200	300	RT	100	RT	RT
a [Å]	18.5597(8)	18.9387(15)	19.3238(12)	18.5303(5)	18.9730(6)	19.3409(16)	19.287(3)	18.5673(9)	19.2235(10)	19.1399(10)
b [Å]	11.9188(3)	11.9783(8)	12.0529(5)	11.8902(4)	11.9295(4)	12.0279(11)	12.0317(17)	11.8735(5)	11.9943(6)	11.8849(6)
c [Å]	11.7871(4)	11.8153(9)	11.8820(6)	11.7514(3)	11.7754(4)	11.8749(12)	11.8591(19)	11.7096(5)	11.8244(6)	11.7325(6)
β [°]	90	90	90	90	90	90	90	90	90	90
V [Å ³]	2607.42(16)	2680.3(3)	2767.4(3)	2589.17(13)	2665.23(15)	2762.5(4)	2751.9(7)	2581.5(2)	2726.4(2)	2668.9(2)
Z	4			4				4		4

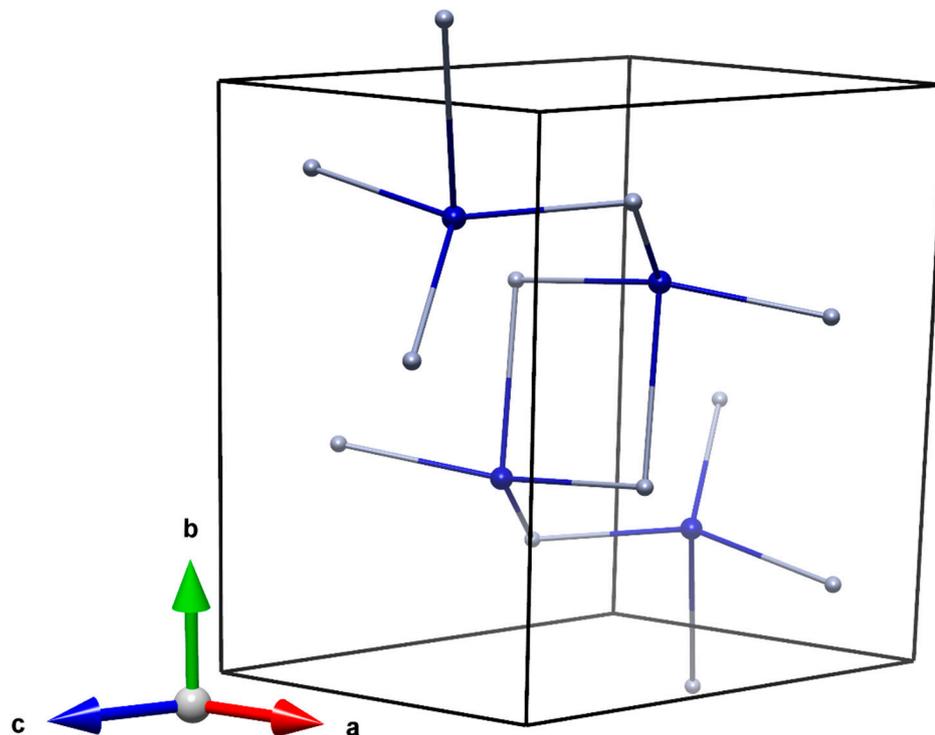


Figure S6. Nitrogen centers of TBA⁺ cations (marked gray) and RE³⁺ (here RE = Tm, marked blue) in α -TBAREB.

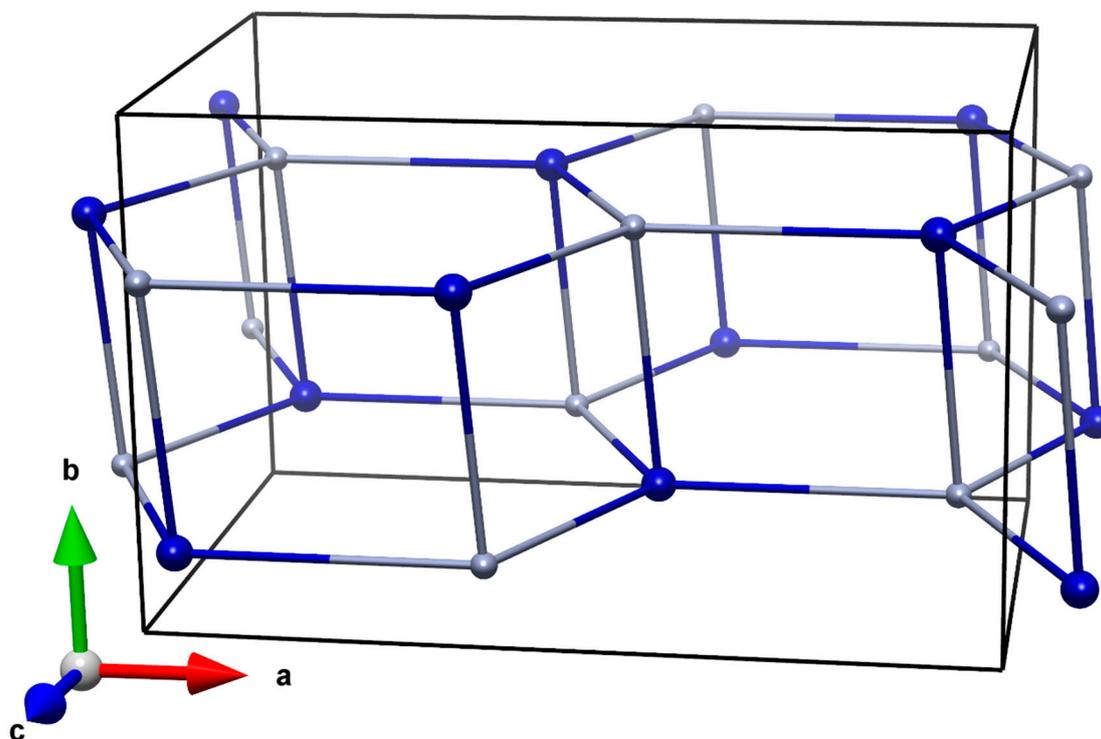


Figure S7. Honeycomb-like structure of nitrogen centers of TBA⁺ cations (marked gray) and RE³⁺ (here RE = Tm, marked blue) in β -TBAREB.

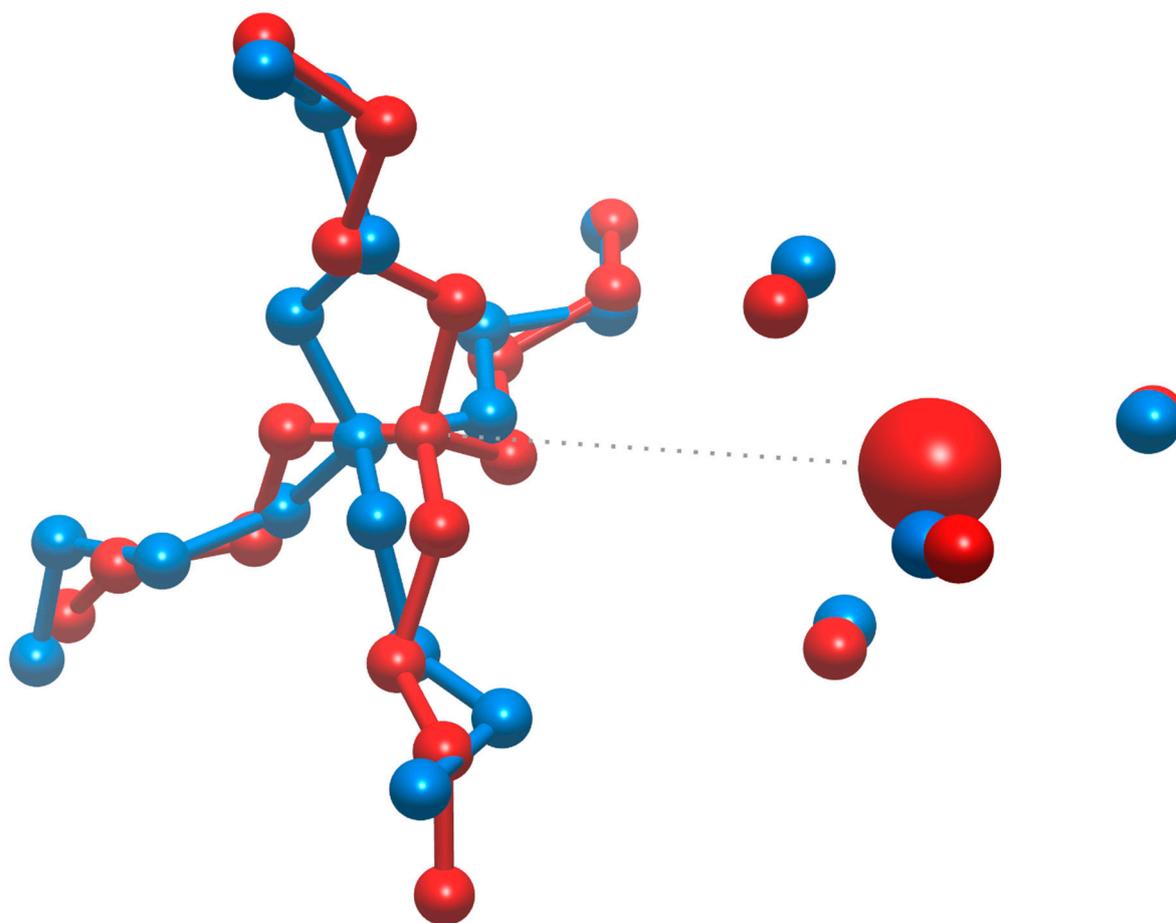


Figure S8. RE...N distances for α -TBAHoB (atoms marked red) and α -TBA_{Tm}B (atoms marked blue), crystal structures obtained from PXRD. Left: TBA⁺, right: [RE(BH₄)₄]⁻. H atoms are not included.

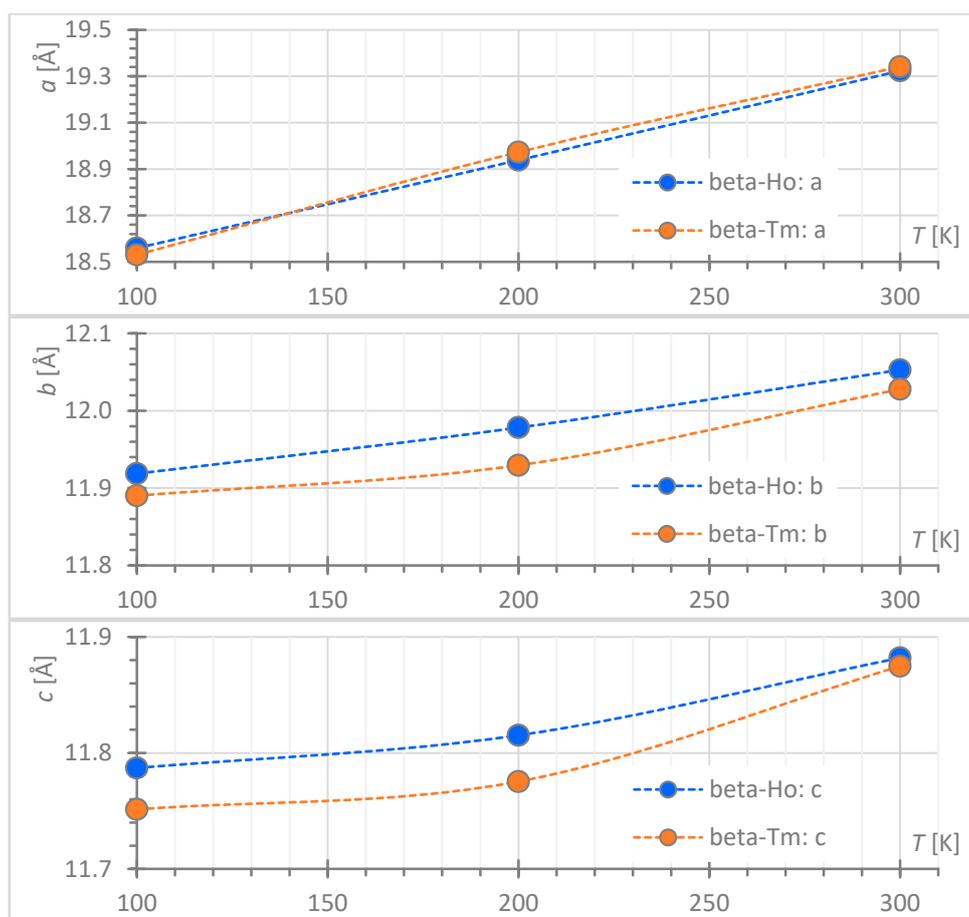


Figure S9. Evolution of a , b , and c lattice parameters of β -TBAHoB and β -TBATmB in the function of temperature. SC-XRD data.

2. Thermal decomposition

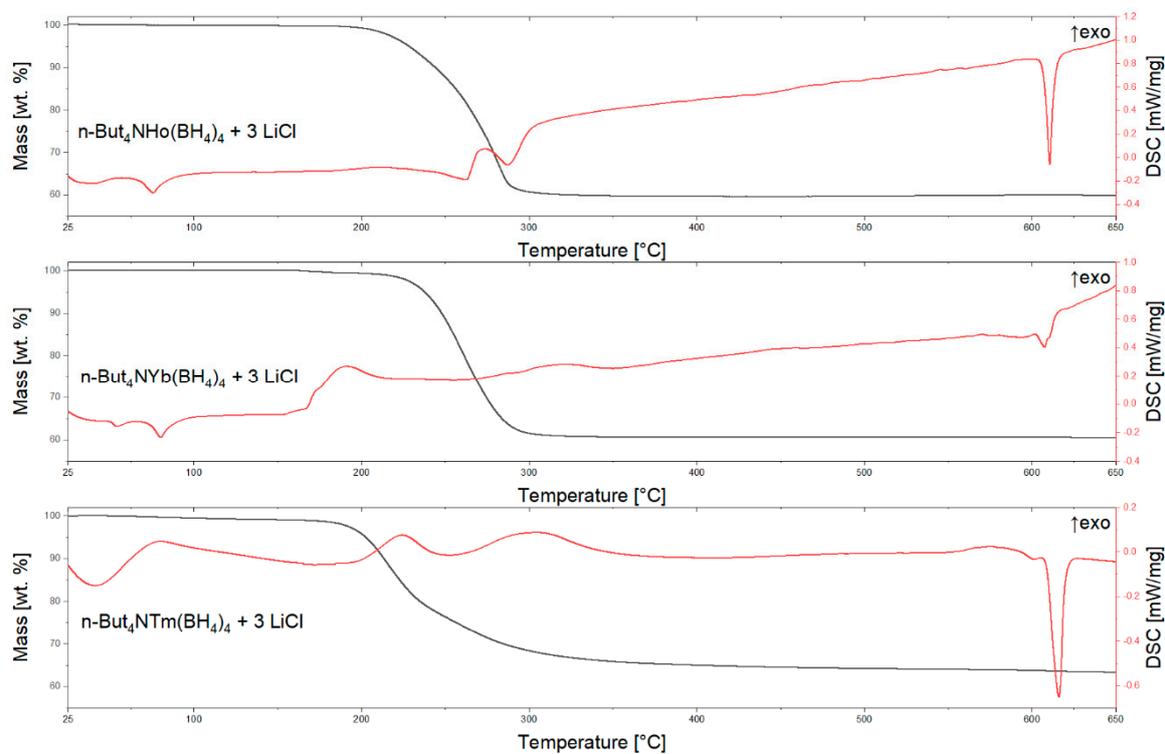


Figure S10. TGA/DSC curves of the as-milled samples (a) **Ho**, (b) **Yb** and (c) **Tm**, up to 650 °C.

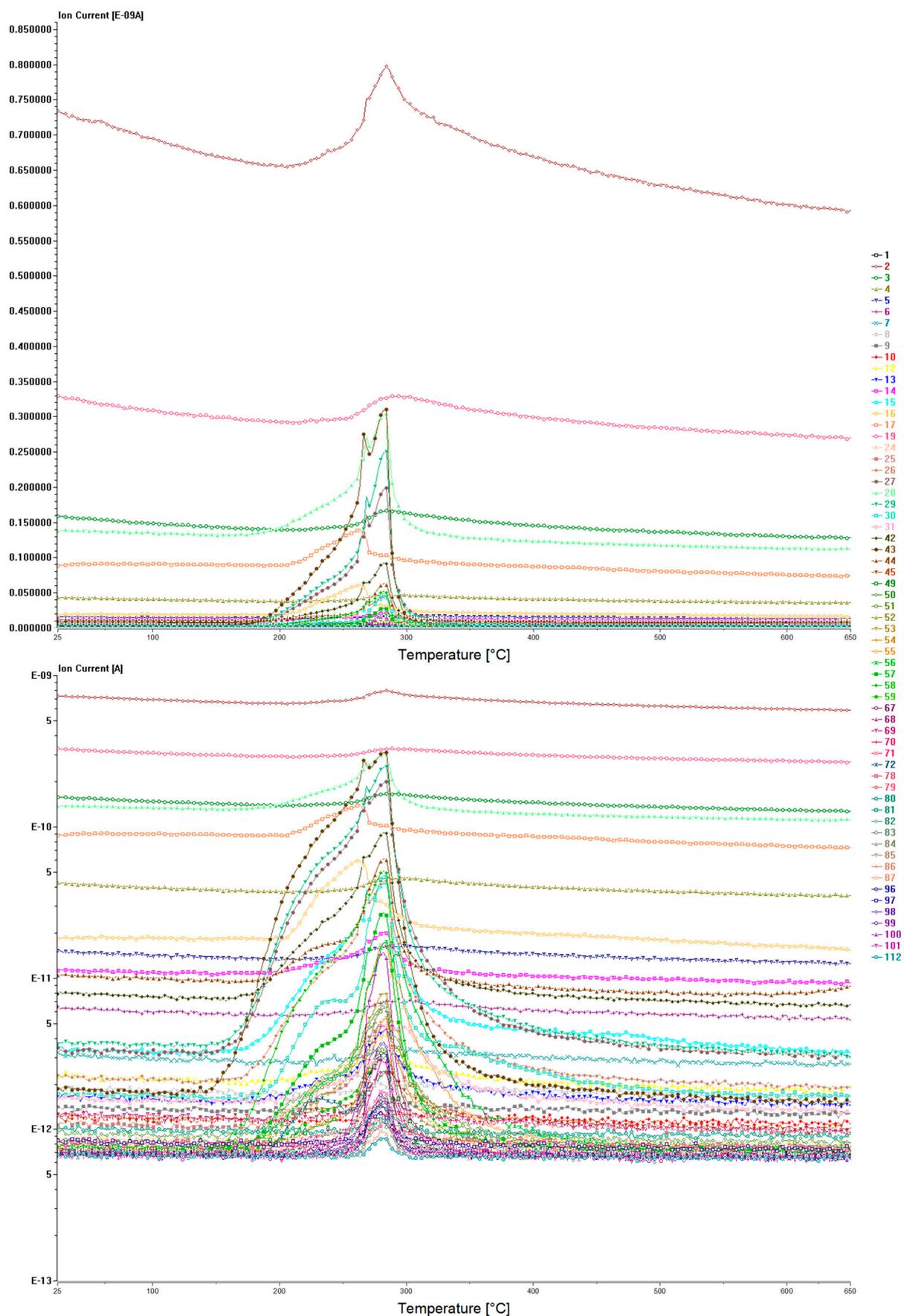


Figure S11. MS spectrum for Ho sample. Top: linear scale; bottom: logarithmic scale; right: m/z value legend.

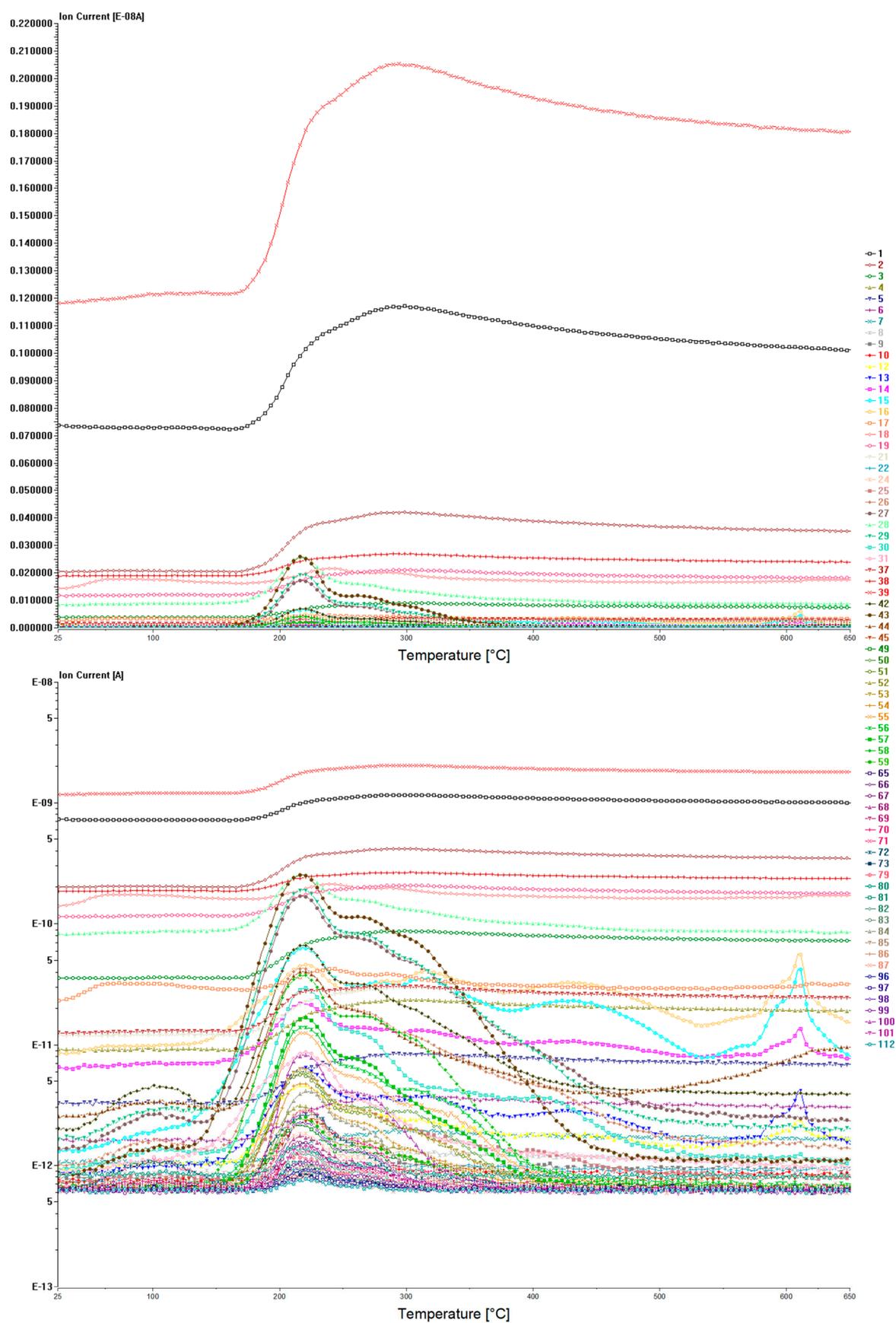


Figure S12. MS spectrum for Tm sample. Top: linear scale; bottom: logarithmic scale; right: m/z value legend.

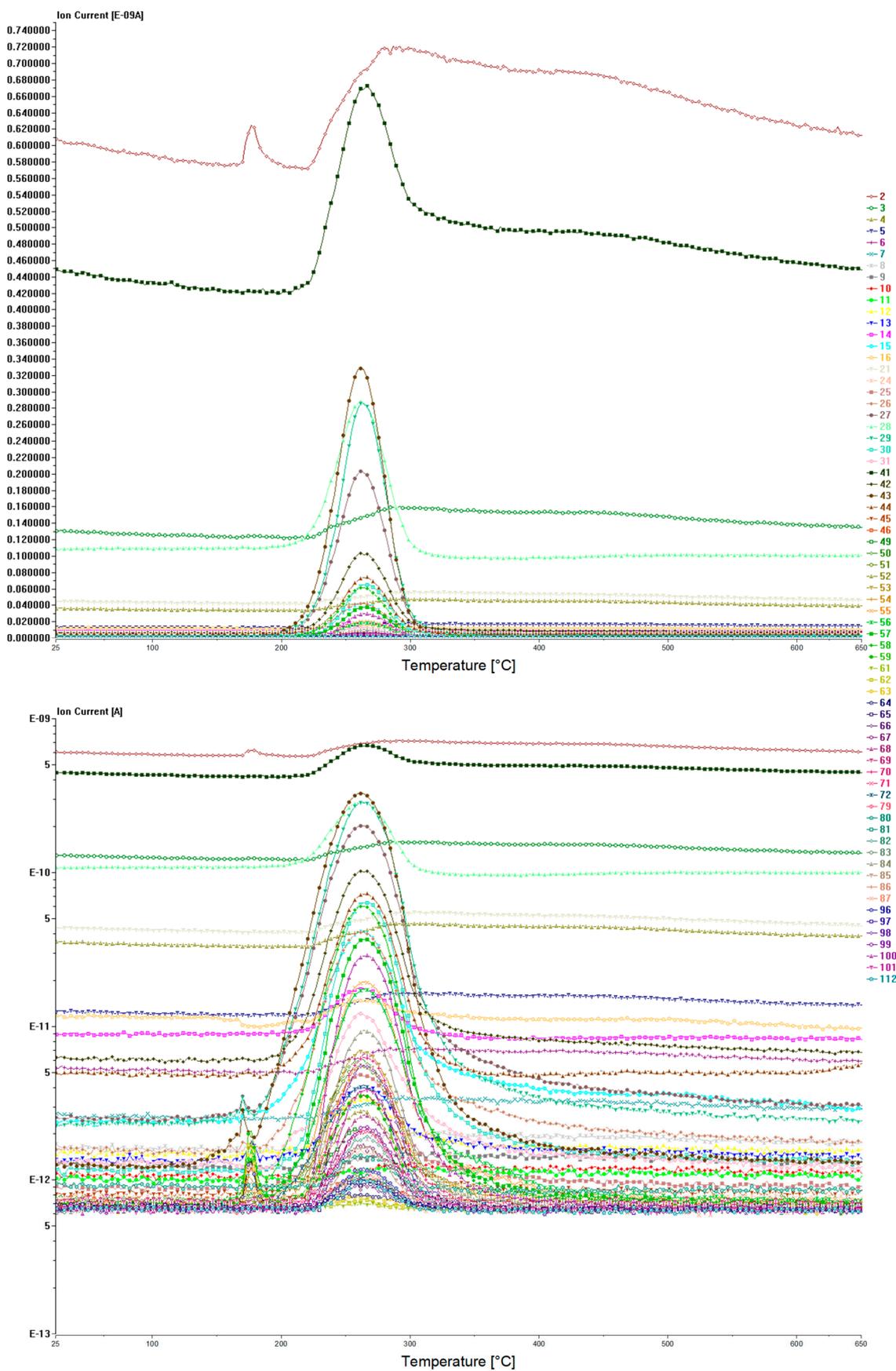


Figure S13. MS spectrum for Yb sample. Top: linear scale; bottom: logarithmic scale; right: m/z value legend.

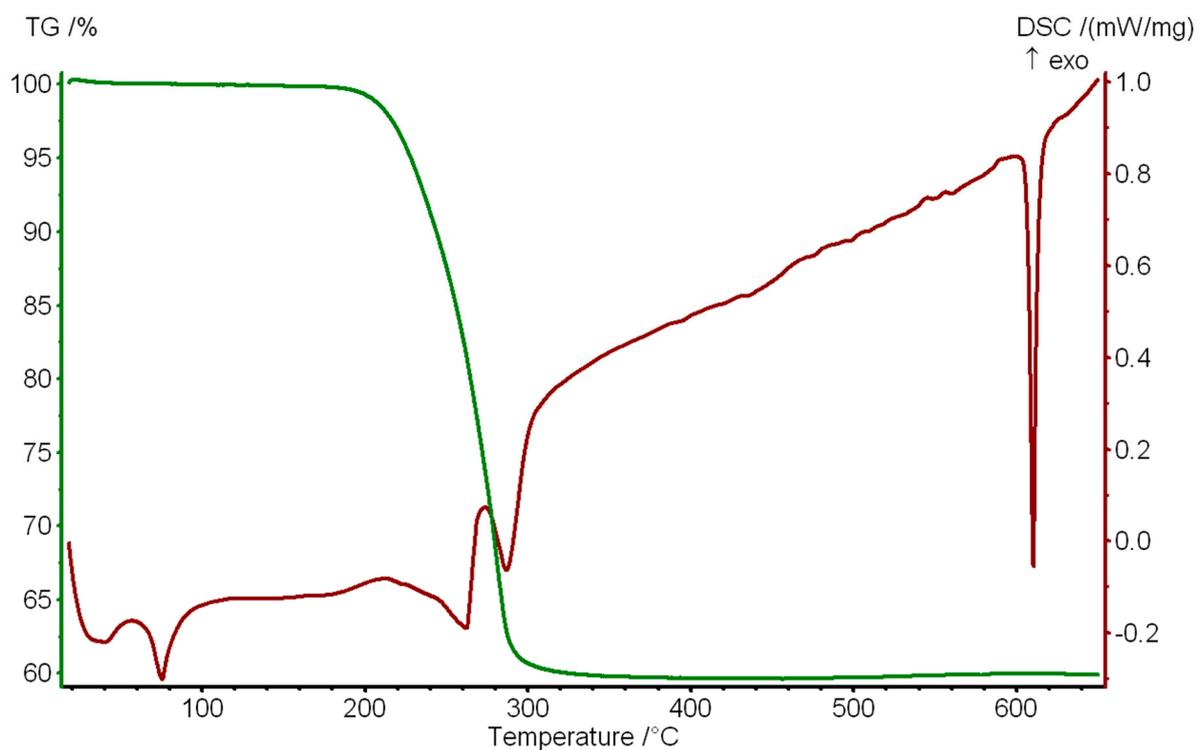


Figure S14. TGA/DSC curves for Ho sample.

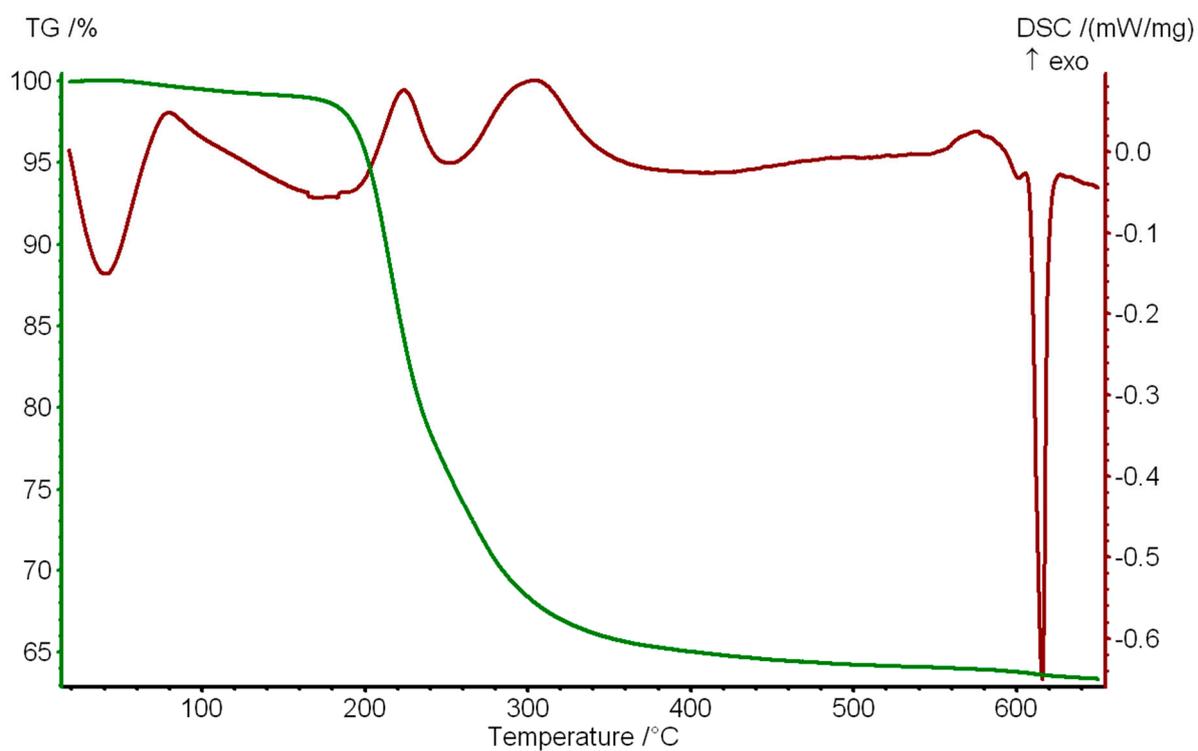


Figure S15. TGA/DSC curves for Tm sample.

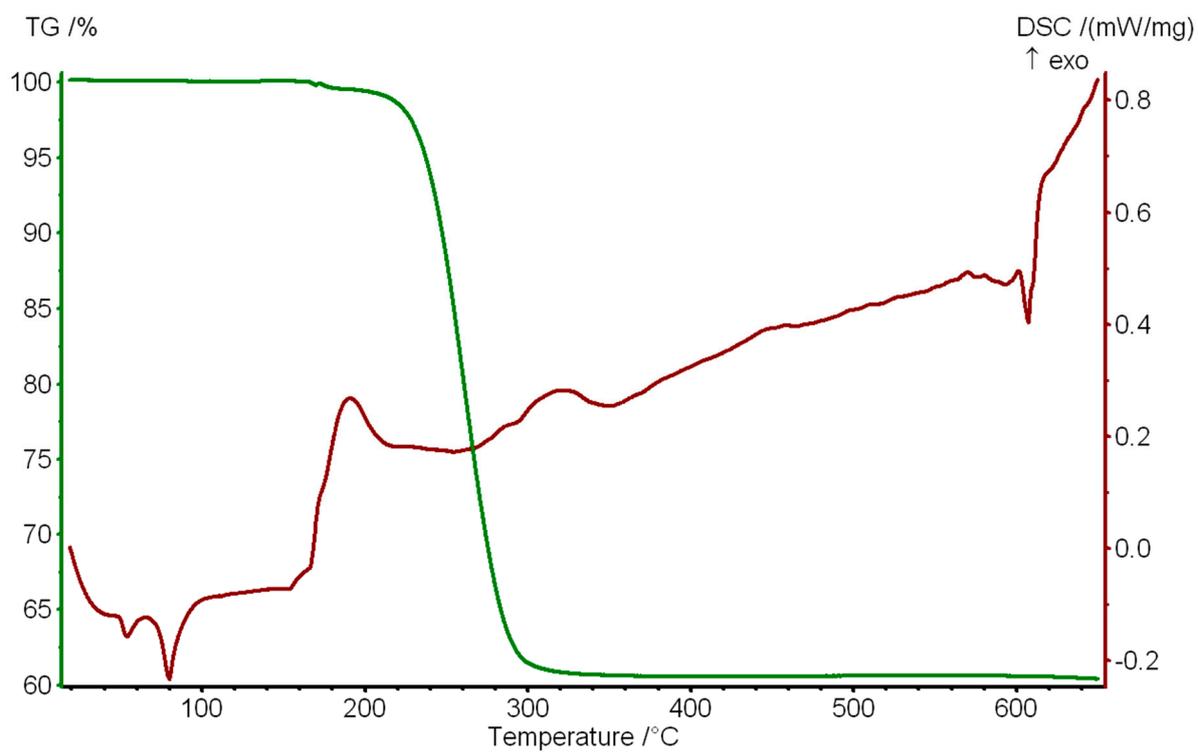


Figure S16. TGA/DSC curves for Yb sample.

3. Solid decomposition products

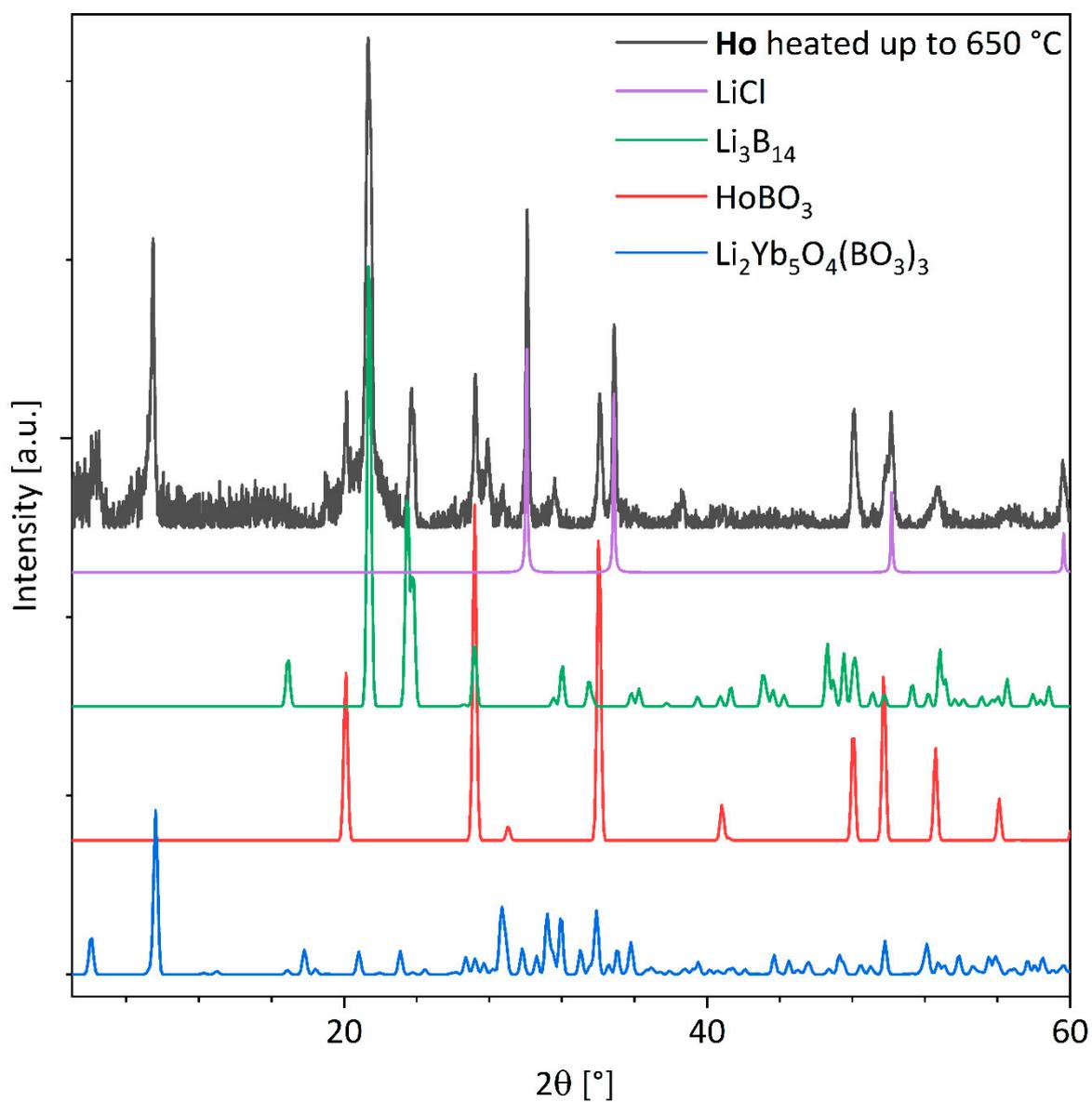


Figure S17. PXRD pattern (with subtracted background) of **Ho** sample heated to 650 °C, and simulated patterns of identified crystalline pyrolysis products [66-69].

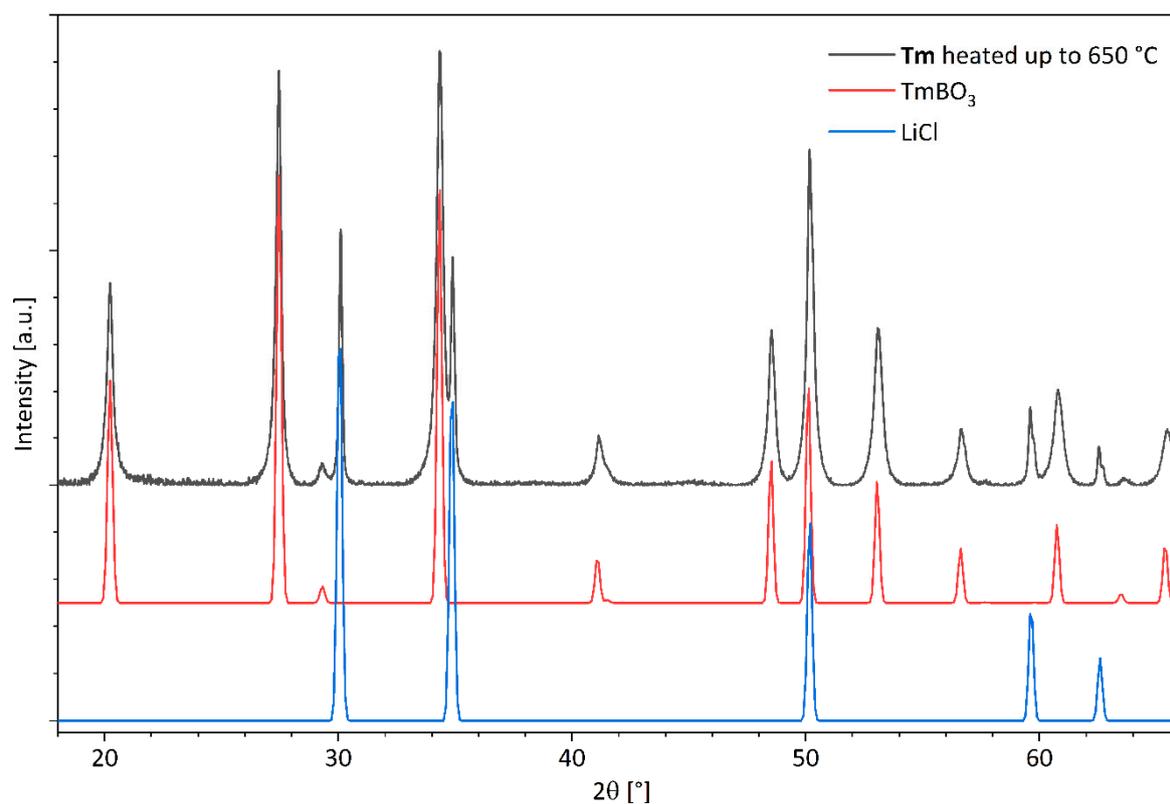


Figure S18. PXRD pattern (with subtracted background) of **Tm** sample heated to 650 °C, and simulated patterns of identified crystalline pyrolysis products [67,69].

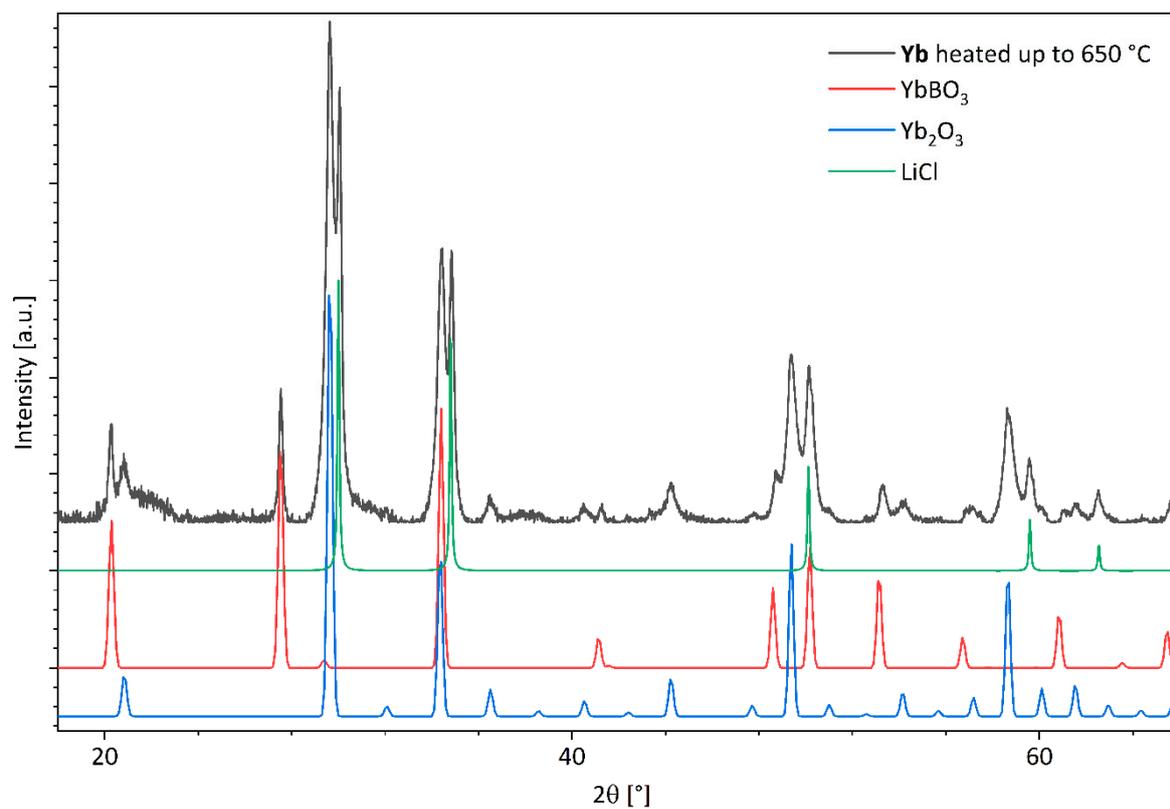


Figure S19. PXRD pattern (with subtracted background) of **Yb** sample heated to 650 °C, and simulated patterns of identified crystalline pyrolysis products [67,69,70].

4. Preliminary CIF of α -TBAHoB from SC-XRD data (100 K)

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Bourhis, L.J., Dolomanov, O.V., Gildea, R.J., Howard, J.A.K., Puschmann, H.

(2015). Acta Cryst. A71, 59-75.

Dolomanov, O.V., Bourhis, L.J., Gildea, R.J., Howard, J.A.K. & Puschmann, H.

(2009), J. Appl. Cryst. 42, 339-341.

Sheldrick, G.M. (2015). Acta Cryst. A71, 3-8.

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using equivalent radius and absorption coefficient.

Empirical absorption correction using spherical harmonics,
implemented in SCALE3 ABSPACK scaling algorithm.

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2	\w	-22.00	71.00	1.00	4.00	--	0.00	38.00	90.00	93
3	\w	-22.00	71.00	1.00	4.00	--	0.00	38.00-180.00		93
4	\w	32.00	106.00	1.00	16.00	--	68.24	-99.00	30.00	74
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6	\w	54.00	112.00	1.00	16.00	--	68.24	-125.00-180.00		58
7	\w	-2.00	94.00	1.00	16.00	--	68.24	-57.00-180.00		96
8	\w	-3.00	90.00	1.00	16.00	--	68.24	-38.00-120.00		93
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_reflns_odcompleteness_iscentric	1
_reflns_odcompleteness_theta	54.18
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_computing_data_collection	'CrysAlisPro 1.171.39.46 (Rigaku OD, 2018)'
_computing_data_reduction	'CrysAlisPro 1.171.39.46 (Rigaku OD, 2018)'
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_computing_publication_material	'Olex2 (Dolomanov et al., 2009)'
_computing_structure_refinement	'olex2.refine (Bourhis et al., 2015)'
_computing_structure_solution	'ShelXT (Sheldrick, 2015)'
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_refine_ls_d_res_low	10.0270
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_refine_ls_hydrogen_treatment	mixed
_refine_ls_matrix_type	full
_refine_ls_number_constraints	53
_refine_ls_number_parameters	252
_refine_ls_number_reflns	3228
_refine_ls_number_restraints	40
_refine_ls_R_factor_all	0.2576
_refine_ls_R_factor_gt	0.1996
_refine_ls_restrained_S_all	1.6352
_refine_ls_shift/su_max	0.0767
_refine_ls_shift/su_mean	0.0038
_refine_ls_structure_factor_coef	Fsqd
_refine_ls_weighting_details	
	'w=1/[\sigma^2(Fo^2)+(0.2P)^2] where P=(Fo^2+2Fc^2)/3'
_refine_ls_weighting_scheme	calc
_refine_ls_wR_factor_gt	0.4724
_refine_ls_wR_factor_ref	0.5140

_olex2_refinement_description

;

1. Fixed Uiso

At 1.2 times of:

All C(H,H) groups

At 1.5 times of:

All C(H,H,H) groups

2. Restrained distances

B00L-H5aa \ \sim B00L-Hl \ \sim B00L-Hq \ \sim B00L-Hp \ \sim B00M-H6aa \ \sim

B00M-Hf \ \sim B00M-Hi ~

B00I-Hh \ \sim B00I-H3aa \ \sim B00I-Hj \ \sim B00I-Hm \ \sim B00M-Hn \ \sim

B00J-Ho \ \sim B00J-H4aa \ \sim B00J-

Hg \ \sim B00J-Hk

with sigma of 0.01

3. Restrained angles

Hk-B00J-Ho

fixed at 109.471 with sigma of 0.02

Hk-B00J-H4aa

fixed at 109.471 with sigma of 0.02

Hk-B00J-Hg

fixed at 109.471 with sigma of 0.02

Ho-B00J-H4aa

fixed at 109.471 with sigma of 0.02

Ho-B00J-Hg

fixed at 109.471 with sigma of 0.02

H4aa-B00J-Hg

fixed at 109.471 with sigma of 0.02

H3aa-B00I-Hj

fixed at 109.471 with sigma of 0.02

H3aa-B00I-Hm

fixed at 109.471 with sigma of 0.02

H3aa-B00I-Hh

fixed at 109.471 with sigma of 0.02

Hj-B00I-Hm

fixed at 109.471 with sigma of 0.02

Hj-B00I-Hh

fixed at 109.471 with sigma of 0.02

Hm-B00I-Hh

fixed at 109.471 with sigma of 0.02

Hp-B00L-Hq

fixed at 109.471 with sigma of 0.02

Hp-B00L-Hl

fixed at 109.471 with sigma of 0.02

Hp-B00L-H5aa

fixed at 109.471 with sigma of 0.02

Hq-B00L-HI

fixed at 109.471 with sigma of 0.02

Hq-B00L-H5aa

fixed at 109.471 with sigma of 0.02

HI-B00L-H5aa

fixed at 109.471 with sigma of 0.02

Hf-B00M-H6aa

fixed at 109.471 with sigma of 0.02

Hf-B00M-Hi

fixed at 109.471 with sigma of 0.02

Hf-B00M-Hn

fixed at 109.471 with sigma of 0.02

H6aa-B00M-Hi

fixed at 109.471 with sigma of 0.02

H6aa-B00M-Hn

fixed at 109.471 with sigma of 0.02

Hi-B00M-Hn

fixed at 109.471 with sigma of 0.02

4. Uiso/Uanisotropy restraints and constraints

Uiso(Hk) = Uiso(H4aa) = Uiso(Ho) = Uiso(Hg) = Uiso(Hp) = Uiso(Hq) = Uiso(HI) =

Uiso(H5aa) = Uiso(H6aa) = Uiso(Hf) = Uiso(Hn) = Uiso(Hi) = Uiso(Hh) =

Uiso(H3aa) = Uiso(Hj) = Uiso(Hm)

5.a Secondary CH2 refined with riding coordinates:

C003(H00a,H00b), C004(H00c,H00d), C005(H00e,H00f), C006(H00g,H00h), C007(H00i,
H00j), C008(H00k,H00l), C00A(H00p,H00q), C00B(H00r,H00s), C00E(H00z,H),
C00F(H00,Ha), C00G(H0aa,Hb), C00H(H1aa,Hc)

5.b Idealised Me refined as rotating group:

C009(H00m,H00n,H00o), C00C(H00t,H00u,H00v), C00D(H00w,H00x,H00y), C00K(H2aa,
Hd,He)

;

_atom_sites_solution_primary dual

loop_

_atom_site_label

_atom_site_type_symbol

_atom_site_fract_x

_atom_site_fract_y

_atom_site_fract_z

_atom_site_U_iso_or_equiv

_atom_site_adp_type

_atom_site_occupancy

_atom_site_refinement_flags_posn

Ho01 Ho 0.79220(19) 0.68774(10) 0.46389(19) 0.0662(11) Uani 1.000000 .
N002 N 0.2623(18) 0.6236(11) 0.3505(17) 0.053(6) Uani 1.000000 .
C003 C 0.279(2) 0.7493(14) 0.374(3) 0.059(7) Uani 1.000000 .
H00a H 0.327(2) 0.7455(14) 0.450(3) 0.071(9) Uiso 1.000000 R
H00b H 0.335(2) 0.7504(14) 0.319(3) 0.071(9) Uiso 1.000000 R
C004 C 0.326(3) 0.6174(16) 0.248(2) 0.068(8) Uani 1.000000 .
H00c H 0.267(3) 0.6279(16) 0.180(2) 0.082(10) Uiso 1.000000 R
H00d H 0.389(3) 0.6513(16) 0.255(2) 0.082(10) Uiso 1.000000 R
C005 C 0.384(3) 0.5513(16) 0.227(3) 0.078(10) Uani 1.000000 .
H00e H 0.446(3) 0.5406(16) 0.293(3) 0.094(12) Uiso 1.000000 R
H00f H 0.322(3) 0.5166(16) 0.220(3) 0.094(12) Uiso 1.000000 R
C006 C 0.446(3) 0.5525(16) 0.115(2) 0.074(9) Uani 1.000000 .
H00g H 0.470(3) 0.5073(16) 0.100(2) 0.088(11) Uiso 1.000000 R
H00h H 0.383(3) 0.5653(16) 0.051(2) 0.088(11) Uiso 1.000000 R
C007 C 0.079(2) 0.5583(16) 0.232(2) 0.065(8) Uani 1.000000 .
H00i H 0.032(2) 0.5992(16) 0.214(2) 0.078(9) Uiso 1.000000 R
H00j H 0.124(2) 0.5492(16) 0.170(2) 0.078(9) Uiso 1.000000 R
C008 C 0.198(2) 0.6894(12) 0.3501(19) 0.045(6) Uani 1.000000 .
H00k H 0.147(2) 0.6951(12) 0.2754(19) 0.054(7) Uiso 1.000000 R
H00l H 0.144(2) 0.6878(12) 0.4074(19) 0.054(7) Uiso 1.000000 R
C009 C -0.091(3) 0.4830(15) 0.138(3) 0.080(10) Uani 1.000000 .
H00m H -0.045(4) 0.463(10) 0.085(8) 0.120(14) Uiso 1.000000 GR
H00n H -0.152(13) 0.452(9) 0.156(4) 0.120(14) Uiso 1.000000 GR
H00o H -0.131(16) 0.523(2) 0.104(11) 0.120(14) Uiso 1.000000 GR
C00A C 0.207(3) 0.8140(13) 0.365(3) 0.063(8) Uani 1.000000 .
H00p H 0.161(3) 0.8181(13) 0.287(3) 0.075(9) Uiso 1.000000 R
H00q H 0.147(3) 0.8113(13) 0.417(3) 0.075(9) Uiso 1.000000 R
C00B C 0.169(2) 0.5660(15) 0.349(2) 0.064(8) Uani 1.000000 .
H00r H 0.213(2) 0.5244(15) 0.365(2) 0.077(9) Uiso 1.000000 R
H00s H 0.121(2) 0.5733(15) 0.410(2) 0.077(9) Uiso 1.000000 R
C00C C 0.283(3) 0.8772(16) 0.394(3) 0.093(11) Uani 1.000000 .
H00t H 0.335(17) 0.884(7) 0.337(12) 0.139(17) Uiso 1.000000 GR
H00u H 0.229(3) 0.915(3) 0.39(2) 0.139(17) Uiso 1.000000 GR
H00v H 0.333(18) 0.873(5) 0.469(10) 0.139(17) Uiso 1.000000 GR
C00D C 0.552(2) 0.5963(14) 0.114(2) 0.064(8) Uani 1.000000 .
H00w H 0.558(10) 0.608(8) 0.036(3) 0.096(12) Uiso 1.000000 GR
H00x H 0.543(8) 0.636(4) 0.157(13) 0.096(12) Uiso 1.000000 GR
H00y H 0.626(3) 0.573(3) 0.149(14) 0.096(12) Uiso 1.000000 GR
C00E C 0.310(2) 0.6162(13) 0.572(2) 0.052(7) Uani 1.000000 .
H00z H 0.259(2) 0.5773(13) 0.578(2) 0.063(8) Uiso 1.000000 R

H H 0.261(2) 0.6559(13) 0.577(2) 0.063(8) Uiso 1.000000 R
C00F C -0.007(3) 0.5009(15) 0.246(3) 0.078(9) Uani 1.000000 .
H00 H 0.042(3) 0.4620(15) 0.273(3) 0.093(11) Uiso 1.000000 R
Ha H -0.056(3) 0.5125(15) 0.305(3) 0.093(11) Uiso 1.000000 R
C00G C 0.420(3) 0.6158(17) 0.669(2) 0.077(9) Uani 1.000000 .
H0aa H 0.476(3) 0.5805(17) 0.655(2) 0.092(11) Uiso 1.000000 R
Hb H 0.463(3) 0.6581(17) 0.670(2) 0.092(11) Uiso 1.000000 R
C00H C 0.358(2) 0.6153(15) 0.463(2) 0.062(8) Uani 1.000000 .
H1aa H 0.401(2) 0.5733(15) 0.459(2) 0.074(9) Uiso 1.000000 R
Hc H 0.418(2) 0.6508(15) 0.466(2) 0.074(9) Uiso 1.000000 R
B00I B 0.654(3) 0.7768(17) 0.508(3) 0.066(9) Uani 1.000000 D
B00J B 0.685(2) 0.5817(15) 0.467(3) 0.053(8) Uani 1.000000 D
C00K C 0.378(3) 0.6048(17) 0.783(2) 0.075(9) Uani 1.000000 .
H2aa H 0.349(18) 0.560(3) 0.787(8) 0.113(14) Uiso 1.000000 GR
Hd H 0.446(5) 0.612(10) 0.844(2) 0.113(14) Uiso 1.000000 GR
He H 0.313(13) 0.635(7) 0.790(8) 0.113(14) Uiso 1.000000 GR
B00L B 0.982(2) 0.6864(18) 0.615(3) 0.075(11) Uani 1.000000 D
B00M B 0.849(3) 0.701(2) 0.273(4) 0.094(14) Uani 1.000000 D
H4aa H 0.796(7) 0.614(7) 0.473(13) 0.069(19) Uiso 1.000000 D
H6aa H 0.952(8) 0.738(6) 0.315(12) 0.069(19) Uiso 1.000000 D
Hf H 0.828(13) 0.656(6) 0.356(9) 0.069(19) Uiso 1.000000 D
H3aa H 0.704(13) 0.732(5) 0.592(9) 0.069(19) Uiso 1.000000 D
Hg H 0.654(13) 0.579(7) 0.575(6) 0.069(19) Uiso 1.000000 D
Hh H 0.726(11) 0.780(8) 0.424(9) 0.069(19) Uiso 1.000000 D
Hi H 0.866(13) 0.668(7) 0.175(8) 0.069(19) Uiso 1.000000 D
Hj H 0.537(6) 0.756(7) 0.461(12) 0.069(19) Uiso 1.000000 D
Hk H 0.695(14) 0.518(3) 0.426(12) 0.069(19) Uiso 1.000000 D
H5aa H 0.966(13) 0.744(4) 0.552(10) 0.069(19) Uiso 1.000000 D
Hl H 1.010(13) 0.635(5) 0.547(9) 0.069(19) Uiso 1.000000 D
Hm H 0.649(14) 0.839(3) 0.557(11) 0.069(19) Uiso 1.000000 D
Hn H 0.749(8) 0.743(6) 0.248(13) 0.069(19) Uiso 1.000000 D
Ho H 0.596(10) 0.616(6) 0.394(10) 0.069(19) Uiso 1.000000 D
Hp H 0.875(8) 0.672(7) 0.656(12) 0.069(19) Uiso 1.000000 D
Hq H 1.076(9) 0.695(7) 0.707(9) 0.069(19) Uiso 1.000000 D

loop_

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_atom_site_aniso_U_22
_atom_site_aniso_U_33
_atom_site_aniso_U_12
_atom_site_aniso_U_13

_atom_site_aniso_U_23

Ho01 0.0520(13) 0.0827(19) 0.0601(16) -0.0003(11) -0.0017(10) -0.0014(11)
N002 0.044(12) 0.073(16) 0.040(13) -0.004(10) 0.001(10) 0.003(10)
C003 0.045(15) 0.08(2) 0.056(18) -0.002(14) 0.028(13) -0.006(14)
C004 0.062(18) 0.10(3) 0.035(15) -0.012(16) -0.006(13) 0.006(14)
C005 0.065(19) 0.09(2) 0.06(2) 0.014(17) -0.026(16) -0.001(16)
C006 0.08(2) 0.10(2) 0.030(15) 0.003(17) -0.029(14) 0.012(14)
C007 0.058(17) 0.10(2) 0.031(14) -0.004(16) -0.001(12) 0.011(14)
C008 0.033(12) 0.070(17) 0.026(13) 0.008(12) -0.009(9) -0.012(11)
C009 0.058(18) 0.09(2) 0.09(2) -0.002(15) -0.006(17) 0.016(17)
C00A 0.069(19) 0.057(18) 0.065(19) -0.000(15) 0.019(15) -0.007(14)
C00B 0.053(16) 0.09(2) 0.049(17) -0.008(15) -0.003(13) -0.002(14)
C00C 0.10(3) 0.09(2) 0.10(3) 0.04(2) 0.01(2) 0.01(2)
C00D 0.039(14) 0.10(2) 0.056(18) 0.001(14) 0.012(13) -0.006(15)
C00E 0.031(12) 0.059(17) 0.063(17) 0.002(11) -0.004(12) 0.000(13)
C00F 0.050(17) 0.08(2) 0.09(2) 0.004(15) -0.003(16) -0.002(18)
C00G 0.063(18) 0.12(3) 0.042(17) 0.003(18) 0.009(14) -0.008(16)
C00H 0.042(15) 0.10(2) 0.045(16) 0.009(14) 0.008(12) 0.001(14)
B00I 0.08(2) 0.08(3) 0.034(18) 0.012(19) 0.007(16) -0.005(15)
B00J 0.036(15) 0.08(2) 0.050(19) 0.003(14) 0.012(13) 0.001(15)
C00K 0.061(18) 0.12(3) 0.037(16) -0.004(17) -0.000(13) 0.008(15)
B00L 0.025(14) 0.13(3) 0.05(2) 0.011(16) -0.040(14) 0.001(19)
B00M 0.036(18) 0.12(4) 0.14(4) 0.002(18) 0.04(2) 0.02(3)

5. FTIR spectra of as-milled samples.

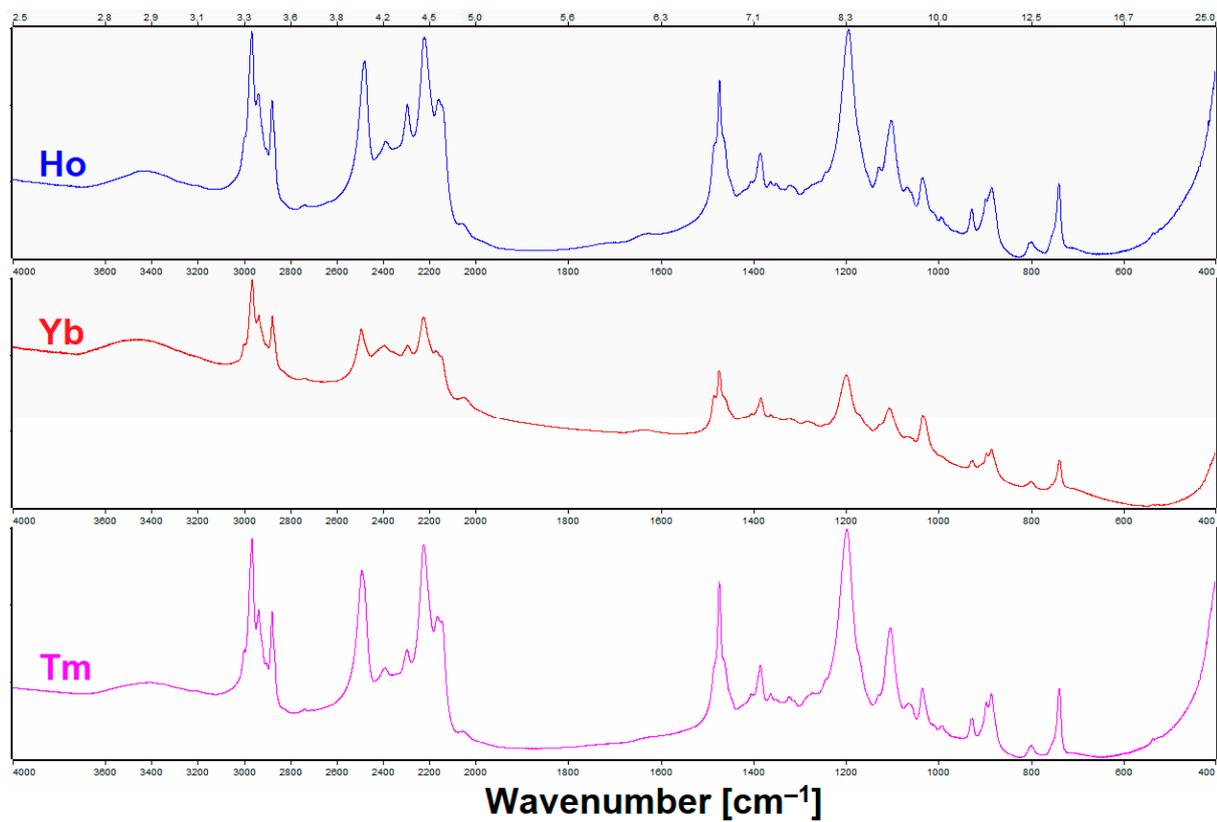


Figure S20. FTIR spectra of Ho, Yb and Tm as-milled samples.